CENTRAL FAX CENTER

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AMENDMENTS TO THE CLAIMS:

Claim 53 is amended. Claims 40 and 41 are cancelled. The following is the status of the claims of the above-captioned application, as amended.

Claims 1-30 (Cancelled.)

- Claim 31. (Currently amended.) An isolated nucleic acid sequence encoding a polypeptide having glucanotransferase activity, wherein the nucleic acid sequence comprises a nucleic acid sequence selected from the group consisting of:
 - (a) a nucleic acid sequence encoding a polypeptide having an amino acid sequence which has at least 80% identity with the amino acid sequence shown as amino acids 1 to 501 of SEQ ID NO:2;
 - a nucleic acid sequence having at least 80% identity with the nucleic acid (b) sequence shown as nucleotides 1 to 1503 of SEQ ID NO:1;
 - a-nucleic-acid-sequence which hybridizes under medium-stringency-conditions-(6) with a complementary-strand of the nucleic acid sequence shown as nucleotides-1 to 1503 of SEQ ID NO:1;
 - (dc) the glucanotransferase encoding part of the DNA sequence cloned into a plasmid present in Escherichia coli DSM 13049; and
 - (<u>ed</u>) a nucleic acid sequence having at least 80% identity to the glucanotransferase encoding part of the DNA sequence cloned into a plasmid present in Escherichia coli DSM 13049.
- Claim 32. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence encoding a polypeptide having an amino acid sequence which has at least 80% identity with the amino acid sequence shown as amino acids 1 to 501 of SEQ ID NO:2.

- Claim 33. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 80% identity with the nucleic acid sequence shown as nucleotides 1 to 1503 of SEQ ID NO:1.
- Claim 34. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 85% identity with the nucleic acid sequence shown as nucleotides 1 to 1503 of SEQ ID NO:1.
- Claim 35. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 90% identity with the nucleic acid sequence shown as nucleotides 1 to 1503 of SEQ ID NO:1.
- Claim 36. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 95% identity with the nucleic acid sequence shown as nucleotides 1 to 1503 of SEQ ID NO:1.
- Claim 37. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 97% identity with the nucleic acid sequence shown as nucleotides 1 to 1503 of SEQ ID NO:1.
- Claim 38. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 98% identity with the nucleic acid sequence shown as nucleotides 1 to 1503 of SEQ ID NO:1.
- Claim 39. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 99% identity with the nucleic acid sequence shown as nucleotides 1 to 1503 of SEQ ID NO:1.
- Claim 40. (Canceled.)
- Claim 41. (Canceled.)
- Claim 42. (Previously presented.) The nucleic acid sequence of claim 31, wherein the

- Claim 43. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 80% identity to the DNA sequence cloned into a plasmid present in *Escherichia coli* DSM 13049.
- Claim 44. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 85% identity to the DNA sequence cloned into a plasmid present in *Escherichia coli* DSM 13049.
- Claim 45. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 90% identity to the DNA sequence cloned into a plasmid present in *Escherichia coli* DSM 13049.
- Claim 46. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 95% identity to the DNA sequence cloned into a plasmid present in *Escherichia coli* DSM 13049.
- Claim 47. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 96% identity to the DNA sequence cloned into a plasmid present in *Escherichia coli* DSM 13049.
- Claim 48. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 97% identity to the DNA sequence cloned into a plasmid present in *Escherichia coli* DSM 13049.
- Claim 49. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 98% identity to the DNA sequence cloned into a plasmid present in *Escherichia coli* DSM 13049.
- Claim 50. (Previously presented.) The nucleic acid sequence of claim 31, wherein the nucleic acid sequence comprises a nucleic acid sequence having at least 99% identity to the

DNA sequence cloned into a plasmid present in Escherichia coli DSM 13049.

- Claim 51. (Previously presented.) A nucleic acid construct comprising the nucleic acid sequence of claim 31 operably linked to one or more control sequences capable of directing the expression of the polypeptide in a suitable expression host.
- Claim 52. (Previously presented.) A recombinant expression vector comprising the nucleic acid construct of claim 51, a promoter, and transcriptional and translational stop signals.
- Claim 53. (Currently amended.) An isolated recombinant host cell comprising the nucleic acid construct of claim 51.
- Claim 54. (Previously presented.) A method for producing polypeptide having glucanotransferase activity, the method comprising:
 - (a) cultivating a recombinant host cell as defined in claim 53 under conditions conducive to the production of the polypeptide; and
 - (b) recovering the polypeptide.